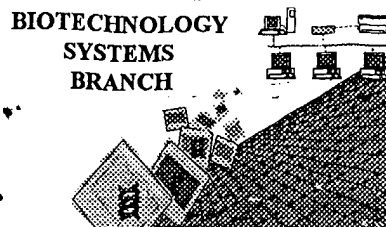


RAW SEQUENCE LISTING ERROR REPORT



The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number: 10/615,479

Source: IPWO

Date Processed by STIC: 12/10/03

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.

PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:

- 1) INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANT, WITH A NOTICE TO COMPLY or,
- 2) TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY

FOR CRF SUBMISSION AND PATENTIN SOFTWARE QUESTIONS, PLEASE CONTACT MARK SPENCER, TELEPHONE: 703-308-4212; FAX: 703-308-4221

Effective 12/13/03: TELEPHONE: 571-272-2510; FAX: 571-273-0221

TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE CHECKER VERSION 4.1 PROGRAM, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW FOR ADDRESS:

<http://www.uspto.gov/web/offices/pac/checker/chkr41note.htm>

Applicants submitting genetic sequence information electronically on diskette or CD-Rom should be aware that there is a possibility that the disk/CD-Rom may have been affected by treatment given to all incoming mail.

Please consider using alternate methods of submission for the disk/CD-Rom or replacement disk/CD-Rom.

Any reply including a sequence listing in electronic form should NOT be sent to the 20231 zip code address for the United States Patent and Trademark Office, and instead should be sent via the following to the indicated addresses:

1. EFS-Bio (<<http://www.uspto.gov/ebc/efs/downloads/documents.htm>> , EFS Submission User Manual - ePAVE)
2. U.S. Postal Service: Commissioner for Patents, P.O. Box 1450, Alexandria, VA 22313-1450
3. Hand Carry directly to (EFFECTIVE 12/01/03):
U.S. Patent and Trademark Office, Box Sequence, Customer Window, Lobby, Room 1B03, Crystal Plaza Two, 2011 South Clark Place, Arlington, VA 22202
4. Federal Express, United Parcel Service, or other delivery service to: U.S. Patent and Trademark Office, Box Sequence, Room 1B03-Mailroom, Crystal Plaza Two, 2011 South Clark Place, Arlington, VA 22202

Revised 10/08/03

Raw Sequence Listing Error Summary

ERROR DETECTED

SUGGESTED CORRECTION

SERIAL NUMBER: 10/615,479

ATTN: NEW RULES CASES: PLEASE DISREGARD ENGLISH "ALPHA" HEADERS, WHICH WERE INSERTED BY PTO SOFTWARE

- 1 ☐ Wrapped Nucleics
Wrapped Aminos The number/text at the end of each line "wrapped" down to the next line. This may occur if your file was retrieved in a word processor after creating it. Please adjust your right margin to .3; this will prevent "wrapping."
- 2 ☐ Invalid Line Length The rules require that a line not exceed 72 characters in length. This includes white spaces.
- 3 ☒ Misaligned Amino
Numbering The numbering under each 5th amino acid is misaligned. Do not use tab codes between numbers; use space characters, instead.
- 4 ☐ Non-ASCII The submitted file was not saved in ASCII(DOS) text, as required by the Sequence Rules. Please ensure your subsequent submission is saved in ASCII text.
- 5 ☐ Variable Length Sequence(s) contain n's or Xaa's representing more than one residue. Per Sequence Rules, each n or Xaa can only represent a single residue. Please present the maximum number of each residue having variable length and indicate in the <220>-<223> section that some may be missing.
- 6 ☐ PatentIn 2.0
"bug" A "bug" in PatentIn version 2.0 has caused the <220>-<223> section to be missing from amino acid sequences(s) . Normally, PatentIn would automatically generate this section from the previously coded nucleic acid sequence. Please manually copy the relevant <220>-<223> section to the subsequent amino acid sequence. This applies to the mandatory <220>-<223> sections for Artificial or Unknown sequences.
- 7 ☐ Skipped Sequences
(OLD RULES) Sequence(s) missing. If intentional, please insert the following lines for each skipped sequence:
(2) INFORMATION FOR SEQ ID NO:X: (insert SEQ ID NO where "X" is shown)
(i) SEQUENCE CHARACTERISTICS: (Do not insert any subheadings under this heading)
(xi) SEQUENCE DESCRIPTION:SEQ ID NO:X: (insert SEQ ID NO where "X" is shown)
This sequence is intentionally skipped

Please also adjust the "(ii) NUMBER OF SEQUENCES:" response to include the skipped sequences.
- 8 ☐ Skipped Sequences
(NEW RULES) Sequence(s) missing. If intentional, please insert the following lines for each skipped sequence.
<210> sequence id number
<400> sequence id number
000
- 9 ☐ Use of n's or Xaa's
(NEW RULES) Use of n's and/or Xaa's have been detected in the Sequence Listing.
Per 1.823 of Sequence Rules, use of <220>-<223> is MANDATORY if n's or Xaa's are present.
In <220> to <223> section, please explain location of n or Xaa, and which residue n or Xaa represents.
- 10 ☐ Invalid <213>
Response Per 1.823 of Sequence Rules, the only valid <213> responses are: Unknown, Artificial Sequence, or scientific name (Genus/species). <220>-<223> section is required when <213> response is Unknown or is Artificial Sequence
- 11 ☐ Use of <220> Sequence(s) missing the <220> "Feature" and associated numeric identifiers and responses.
Use of <220> to <223> is MANDATORY if <213> "Organism" response is "Artificial Sequence" or "Unknown." Please explain source of genetic material in <220> to <223> section.
(See "Federal Register," 00/01/1998, Vol. 63, No. 104, pp. 29631-32) (Sec. 1.823 of Sequence Rules)
- 12 ☐ PatentIn 2.0
"bug" Please do not use "Copy to Disk" function of PatentIn version 2.0. This causes a corrupted file, resulting in missing mandatory numeric identifiers and responses (as indicated on raw sequence listing). Instead, please use "File Manager" or any other manual means to copy file to floppy disk.
- 13 ☐ Misuse of n/Xaa "n" can only represent a single nucleotide; "Xaa" can only represent a single amino acid



IFWO

RAW SEQUENCE LISTING

DATE: 12/10/2003

PATENT APPLICATION: US/10/615,479

TIME: 17:33:46

Input Set : A:\PTO.YF.txt

Output Set: N:\CRF4\12102003\J615479.raw

3 <110> APPLICANT: Larson Mr., Richard S.
 5 <120> TITLE OF INVENTION: Peptide Inhibitors of LFA-1/ICAM-1 Interaction
 7 <130> FILE REFERENCE: N12-012.CIP1
 9 <140> CURRENT APPLICATION NUMBER: 10/615,479
 10 <141> CURRENT FILING DATE: 2003-07-08
 12 <160> NUMBER OF SEQ ID NOS: 33
 14 <170> SOFTWARE: Patent In Ver. 4.1

ERRORED SEQUENCES

Does Not Comply
 Corrected Diskette Needed

p1-5

16 <210> SEQ ID NO: 1
 17 <211> LENGTH: 9
 18 <212> TYPE: PRT
 19 <213> ORGANISM: Artificial Sequence
 21 <220> FEATURE:
 22 <223> OTHER INFORMATION: Description of Artificial Sequence:Synthetic
 24 <400> SEQUENCE: 1
 25 Cys Leu Leu Arg Met Arg Ser Arg Cys
 E--> 26 1 *5* *5*
 28 <210> SEQ ID NO: 2
 29 <211> LENGTH: 9
 30 <212> TYPE: PRT
 31 <213> ORGANISM: Artificial Sequence
 33 <220> FEATURE:
 34 <223> OTHER INFORMATION: Description of Artificial Sequence:Synthetic
 36 <400> SEQUENCE: 2
 37 Cys Leu Leu Arg Met Arg Leu Ile Cys
 E--> 38 1 *5* *5*
 40 <210> SEQ ID NO: 3
 41 <211> LENGTH: 9
 42 <212> TYPE: PRT
 43 <213> ORGANISM: Artificial Sequence
 45 <220> FEATURE:
 46 <223> OTHER INFORMATION: Description of Artificial Sequence:Synthetic
 49 <400> SEQUENCE: 3
 50 Cys Leu Leu Arg Met Arg Ser Ile Cys
 E--> 51 1 *5* *5*
 52 <210> SEQ ID NO: 4
 53 <211> LENGTH: 9
 54 <212> TYPE: PRT
 55 <213> ORGANISM: Artificial Sequence
 57 <220> FEATURE:

9
 Misaligned Amino
 Acid numbers.
 See item
9 *3* on Error
 Summary
 Sheet.

RAW SEQUENCE LISTING

PATENT APPLICATION: US/10/615,479

DATE: 12/10/2003

TIME: 17:33:46

Input Set : A:\PTO.YF.txt

Output Set: N:\CRF4\12102003\J615479.raw

58 <223> OTHER INFORMATION: Description of Artificial Sequence:Synthetic
60 <400> SEQUENCE: 4
61 Cys Leu Leu Arg Met Arg Leu Arg Cys
E--> 62 1 5 5
64 <210> SEQ ID NO: 5
65 <211> LENGTH: 9
66 <212> TYPE: PRT
67 <213> ORGANISM: Artificial Sequence
69 <220> FEATURE:
70 <223> OTHER INFORMATION: Description of Artificial Sequence:Synthetic
72 <400> SEQUENCE: 5
73 Cys Leu Leu Arg Pro Arg Ser Ile Cys
E--> 74 1 5 5
76 <210> SEQ ID NO: 6
77 <211> LENGTH: 9
78 <212> TYPE: PRT
79 <213> ORGANISM: Artificial Sequence
81 <220> FEATURE:
82 <223> OTHER INFORMATION: Description of Artificial Sequence:Synthetic
84 <400> SEQUENCE: 6
85 Cys Leu Leu Arg Pro Arg Ser Arg Cys
E--> 86 1 5 5
88 <210> SEQ ID NO: 7
89 <211> LENGTH: 9
90 <212> TYPE: PRT
92 <213> ORGANISM: Artificial Sequence
94 <220> FEATURE:
95 <223> OTHER INFORMATION: Description of Artificial Sequence:Synthetic
97 <400> SEQUENCE: 7
98 Cys Leu Leu Arg Pro Arg Leu Ile Cys
E--> 99 1 5 5
101 <210> SEQ ID NO: 8
102 <211> LENGTH: 9
103 <212> TYPE: PRT
104 <213> ORGANISM: Artificial Sequence
106 <220> FEATURE:
107 <223> OTHER INFORMATION: Description of Artificial Sequence:Synthetic
109 <400> SEQUENCE: 8
110 Cys Leu Leu Arg Pro Arg Leu Arg Cys
E--> 111 1 5 5
113 <210> SEQ ID NO: 9
114 <211> LENGTH: 9
115 <212> TYPE: PRT
116 <213> ORGANISM: Artificial Sequence
118 <220> FEATURE:
119 <223> OTHER INFORMATION: Description of Artificial Sequence:Synthetic
121 <400> SEQUENCE: 9
122 Cys Leu Leu Arg Met Asn Ser Ile Cys
E--> 123 1 5 5

Same error⁹

Same error⁹

Same error⁹

Same error⁹

Same error⁹

RAW SEQUENCE LISTING

PATENT APPLICATION: US/10/615,479

DATE: 12/10/2003

TIME: 17:33:46

Input Set : A:\PTO.YF.txt

Output Set: N:\CRF4\12102003\J615479.raw

125 <210> SEQ ID NO: 10
126 <211> LENGTH: 9
127 <212> TYPE: PRT
128 <213> ORGANISM: Artificial Sequence
130 <220> FEATURE:
131 <223> OTHER INFORMATION: Description of Artificial Sequence:Synthetic
133 <400> SEQUENCE: 10
134 Cys Leu Leu Arg Met Asn Ser Arg Cys
E--> 135 1 . 5
137 <210> SEQ ID NO: 11
138 <211> LENGTH: 9
139 <212> TYPE: PRT
140 <213> ORGANISM: Artificial Sequence
142 <220> FEATURE:
143 <223> OTHER INFORMATION: Description of Artificial Sequence:Synthetic
145 <400> SEQUENCE: 11
146 Cys Leu Leu Arg Met Asn Leu Ile Cys
E--> 147 1 5
149 <210> SEQ ID NO: 12
150 <211> LENGTH: 9
151 <212> TYPE: PRT
152 <213> ORGANISM: Artificial Sequence
154 <220> FEATURE:
155 <223> OTHER INFORMATION: Description of Artificial Sequence:Synthetic
157 <400> SEQUENCE: 12
158 Cys Leu Leu Arg Met Asn Leu Arg Cys
E--> 159 1 5
161 <210> SEQ ID NO: 13
162 <211> LENGTH: 9
163 <212> TYPE: PRT
164 <213> ORGANISM: Artificial Sequence
166 <220> FEATURE:
167 <223> OTHER INFORMATION: Description of Artificial Sequence:Synthetic
169 <400> SEQUENCE: 13
170 Cys Leu Leu Arg Pro Asn Ser Ile Cys
E--> 171 1 5
173 <210> SEQ ID NO: 14
174 <211> LENGTH: 9
175 <212> TYPE: PRT
176 <213> ORGANISM: Artificial Sequence
178 <220> FEATURE:
179 <223> OTHER INFORMATION: Description of Artificial Sequence:Synthetic
181 <400> SEQUENCE: 14
182 Cys Leu Leu Arg Pro Asn Ser Arg Cys
E--> 183 1 5
185 <210> SEQ ID NO: 15
186 <211> LENGTH: 9
187 <212> TYPE: PRT
188 <213> ORGANISM: Artificial Sequence

Same error 9

Same error 9

Same error 9

Same error 9

RAW SEQUENCE LISTING

PATENT APPLICATION: US/10/615,479

DATE: 12/10/2003

TIME: 17:33:46

Input Set : A:\PTO.YF.txt

Output Set: N:\CRF4\12102003\J615479.raw

190 <220> FEATURE:
191 <223> OTHER INFORMATION: Description of Artificial Sequence:Synthetic.
193 <400> SEQUENCE: 15
194 Cys Leu Leu Arg Pro Asn Leu Ile Cys 9
E--> 195 1 5
197 <210> SEQ ID NO: 16
198 <211> LENGTH: 9
199 <212> TYPE: PRT
200 <213> ORGANISM: Artificial Sequence
202 <220> FEATURE:
203 <223> OTHER INFORMATION: Description of Artificial Sequence:Synthetic
205 <400> SEQUENCE: 16
206 Cys Leu Leu Arg Pro Asn Leu Arg Cys 9
E--> 207 1 5
209 <210> SEQ ID NO: 17
210 <211> LENGTH: 9
211 <212> TYPE: PRT
212 <213> ORGANISM: Artificial Sequence
214 <220> FEATURE:
215 <223> OTHER INFORMATION: Description of Artificial Sequence:Synthetic
217 <400> SEQUENCE: 17
218 Cys Met Leu Arg Met Arg Ser Ile Cys 9
E--> 219 1 5
221 <210> SEQ ID NO: 18
222 <211> LENGTH: 9
223 <212> TYPE: PRT
224 <213> ORGANISM: Artificial Sequence
227 <220> FEATURE:
228 <223> OTHER INFORMATION: Description of Artificial Sequence:Synthetic
230 <400> SEQUENCE: 18
231 Cys Met Leu Arg Met Arg Ser Arg Cys 9
E--> 232 1 5
234 <210> SEQ ID NO: 19
235 <211> LENGTH: 9
236 <212> TYPE: PRT
237 <213> ORGANISM: Artificial Sequence
239 <220> FEATURE:
240 <223> OTHER INFORMATION: Description of Artificial Sequence:Synthetic
242 <400> SEQUENCE: 19
243 Cys Met Leu Arg Met Arg Leu Ile Cys 9
E--> 244 1 5
246 <210> SEQ ID NO: 20
247 <211> LENGTH: 9
248 <212> TYPE: PRT
249 <213> ORGANISM: Artificial Sequence
251 <220> FEATURE:
252 <223> OTHER INFORMATION: Description of Artificial Sequence:Synthetic
254 <400> SEQUENCE: 20
255 Cys Met Leu Arg Met Arg Leu Arg Cys 9

Same error

Same error

Same error

Same error

Same error

RAW SEQUENCE LISTING

PATENT APPLICATION: US/10/615,479

DATE: 12/10/2003

TIME: 17:33:46

Input Set : A:\PTO.YF.txt

Output Set: N:\CRF4\12102003\J615479.raw

E--> 256 1

~~5~~

VERIFICATION SUMMARY

PATENT APPLICATION: US/10/615,479

DATE: 12/10/2003

TIME: 17:33:47

Input Set : A:\PTO.YF.txt

Output Set: N:\CRF4\12102003\J615479.raw

L:26 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:1
L:26 M:301 E: (44) No Sequence Data was Shown, SEQ ID:1
L:26 M:252 E: No. of Seq. differs, <211> LENGTH:Input:9 Found:0 SEQ:1
L:38 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:2
L:38 M:301 E: (44) No Sequence Data was Shown, SEQ ID:2
L:38 M:252 E: No. of Seq. differs, <211> LENGTH:Input:9 Found:0 SEQ:2
L:51 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:3
L:51 M:301 E: (44) No Sequence Data was Shown, SEQ ID:3
L:51 M:252 E: No. of Seq. differs, <211> LENGTH:Input:9 Found:0 SEQ:3
L:62 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:4
L:62 M:301 E: (44) No Sequence Data was Shown, SEQ ID:4
L:62 M:252 E: No. of Seq. differs, <211> LENGTH:Input:9 Found:0 SEQ:4
L:74 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:5
L:74 M:301 E: (44) No Sequence Data was Shown, SEQ ID:5
L:74 M:252 E: No. of Seq. differs, <211> LENGTH:Input:9 Found:0 SEQ:5
L:86 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:6
L:86 M:301 E: (44) No Sequence Data was Shown, SEQ ID:6
L:86 M:252 E: No. of Seq. differs, <211> LENGTH:Input:9 Found:0 SEQ:6
L:99 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:7
L:99 M:301 E: (44) No Sequence Data was Shown, SEQ ID:7
L:99 M:252 E: No. of Seq. differs, <211> LENGTH:Input:9 Found:0 SEQ:7
L:111 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:8
L:111 M:301 E: (44) No Sequence Data was Shown, SEQ ID:8
L:111 M:252 E: No. of Seq. differs, <211> LENGTH:Input:9 Found:0 SEQ:8
L:123 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:9
L:123 M:301 E: (44) No Sequence Data was Shown, SEQ ID:9
L:123 M:252 E: No. of Seq. differs, <211> LENGTH:Input:9 Found:0 SEQ:9
L:135 M:301 E: (44) No Sequence Data was Shown, SEQ ID:10
L:135 M:252 E: No. of Seq. differs, <211> LENGTH:Input:9 Found:0 SEQ:10
L:147 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:11
L:147 M:301 E: (44) No Sequence Data was Shown, SEQ ID:11
L:147 M:252 E: No. of Seq. differs, <211> LENGTH:Input:9 Found:0 SEQ:11
L:159 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:12
L:159 M:301 E: (44) No Sequence Data was Shown, SEQ ID:12
L:159 M:252 E: No. of Seq. differs, <211> LENGTH:Input:9 Found:0 SEQ:12
L:171 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:13
L:171 M:301 E: (44) No Sequence Data was Shown, SEQ ID:13
L:171 M:252 E: No. of Seq. differs, <211> LENGTH:Input:9 Found:0 SEQ:13
L:183 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:14
L:183 M:301 E: (44) No Sequence Data was Shown, SEQ ID:14
L:183 M:252 E: No. of Seq. differs, <211> LENGTH:Input:9 Found:0 SEQ:14
L:195 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:15
L:195 M:301 E: (44) No Sequence Data was Shown, SEQ ID:15
L:195 M:252 E: No. of Seq. differs, <211> LENGTH:Input:9 Found:0 SEQ:15
L:207 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:16
L:207 M:301 E: (44) No Sequence Data was Shown, SEQ ID:16
L:207 M:252 E: No. of Seq. differs, <211> LENGTH:Input:9 Found:0 SEQ:16
L:219 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:17

VERIFICATION SUMMARYPATENT APPLICATION: **US/10/615,479**

DATE: 12/10/2003

TIME: 17:33:47

Input Set : **A:\PTO.YF.txt**Output Set: **N:\CRF4\12102003\J615479.raw**

L:219 M:301 E: (44) No Sequence Data was Shown, SEQ ID:17
L:219 M:252 E: No. of Seq. differs, <211> LENGTH:Input:9 Found:0 SEQ:17
L:232 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:18
L:232 M:301 E: (44) No Sequence Data was Shown, SEQ ID:18
L:232 M:252 E: No. of Seq. differs, <211> LENGTH:Input:9 Found:0 SEQ:18
L:244 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:19
L:244 M:301 E: (44) No Sequence Data was Shown, SEQ ID:19
L:244 M:252 E: No. of Seq. differs, <211> LENGTH:Input:9 Found:0 SEQ:19
L:256 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:20
L:256 M:301 E: (44) No Sequence Data was Shown, SEQ ID:20
L:256 M:252 E: No. of Seq. differs, <211> LENGTH:Input:9 Found:0 SEQ:20
L:268 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:21
L:268 M:301 E: (44) No Sequence Data was Shown, SEQ ID:21
L:268 M:252 E: No. of Seq. differs, <211> LENGTH:Input:9 Found:0 SEQ:21
L:280 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:22
L:280 M:301 E: (44) No Sequence Data was Shown, SEQ ID:22
L:280 M:252 E: No. of Seq. differs, <211> LENGTH:Input:9 Found:0 SEQ:22
L:292 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:23
L:292 M:301 E: (44) No Sequence Data was Shown, SEQ ID:23
L:292 M:252 E: No. of Seq. differs, <211> LENGTH:Input:9 Found:0 SEQ:23
L:304 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:24
L:304 M:301 E: (44) No Sequence Data was Shown, SEQ ID:24
L:304 M:252 E: No. of Seq. differs, <211> LENGTH:Input:9 Found:0 SEQ:24
L:316 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:25
L:316 M:301 E: (44) No Sequence Data was Shown, SEQ ID:25
L:316 M:252 E: No. of Seq. differs, <211> LENGTH:Input:9 Found:0 SEQ:25
L:328 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:26
L:328 M:301 E: (44) No Sequence Data was Shown, SEQ ID:26
L:328 M:252 E: No. of Seq. differs, <211> LENGTH:Input:9 Found:0 SEQ:26
L:340 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:27
L:340 M:301 E: (44) No Sequence Data was Shown, SEQ ID:27
L:340 M:252 E: No. of Seq. differs, <211> LENGTH:Input:9 Found:0 SEQ:27
L:352 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:28
L:352 M:301 E: (44) No Sequence Data was Shown, SEQ ID:28
L:352 M:252 E: No. of Seq. differs, <211> LENGTH:Input:9 Found:0 SEQ:28
L:364 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:29
L:364 M:301 E: (44) No Sequence Data was Shown, SEQ ID:29
L:364 M:252 E: No. of Seq. differs, <211> LENGTH:Input:9 Found:0 SEQ:29
L:376 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:30
L:376 M:301 E: (44) No Sequence Data was Shown, SEQ ID:30
L:376 M:252 E: No. of Seq. differs, <211> LENGTH:Input:9 Found:0 SEQ:30
L:388 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:31
L:388 M:301 E: (44) No Sequence Data was Shown, SEQ ID:31
L:388 M:252 E: No. of Seq. differs, <211> LENGTH:Input:9 Found:0 SEQ:31
L:400 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:32
L:400 M:301 E: (44) No Sequence Data was Shown, SEQ ID:32
L:400 M:252 E: No. of Seq. differs, <211> LENGTH:Input:9 Found:0 SEQ:32
L:414 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:33
L:414 M:301 E: (44) No Sequence Data was Shown, SEQ ID:33

VERIFICATION SUMMARY

PATENT APPLICATION: US/10/615,479

DATE: 12/10/2003

TIME: 17:33:47

Input Set : A:\PTO.YF.txt

Output Set: N:\CRF4\12102003\J615479.raw

L:414 M:252 E: No. of Seq. differs, <211> LENGTH:Input:9 Found:0 SEQ:33